DEET AVILLA CONV

SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: SEIDMAN, CHRISTINE

SEIDMAN, JOHN WATKINS, HUGH

ROSENZWEIG, ANTHONY

- (ii) TITLE OF INVENTION: A METHOD FOR DETECTING
 DISEASE-ASSOCIATED MUTATIONS
- (iii) NUMBER OF SEQUENCES: 10
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: LAHIVE & COCKFIELD
 - (B) STREET: 60 STATE STREET, Suite 510
 - (C) CITY: BOSTON
 - (D) STATE: Massachusetts
 - (E) COUNTRY: U.S.A.
 - (F) ZIP: 02109
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: ASCII
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/989,160
 - (B) FILING DATE: 11-DEC-1993
 - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: HANLEY, ELIZABETH A.
 - (B) REGISTRATION NUMBER: 33,505
 - (C) REFERENCE/DOCKET NUMBER: GWL-111
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (617) 227-7400
 - (B) TELEFAX: (617) 227-5941
- (2) INFORMATION FOR SEO ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 24 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA

Fileu: December 11, 1992

| | (XI) SEQUENCE DESCRIPTION: SEQ ID NO:1: | |
|------|--|----|
| CAAG | GATCGC TACGGCTCCT GGAT | 26 |
| (2) | INFORMATION FOR SEQ ID NO:2: | |
| • | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
|) | (ii) MOLECULE TYPE: CDNA | |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2: | |
| GCGG | ATCCAG GTAGGCAGAC TTGTCAGCCT | 30 |
| (2) | INFORMATION FOR SEQ ID NO:3: | |
| | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| | (ii) MOLECULE TYPE: CDNA | |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3: | |
| ATGC | CAACCC TGCTCTGGAG GCCT | 24 |
| (2) | INFORMATION FOR SEQ ID NO:4: | |
| | (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear | |
| | (ii) MOLECULE TYPE: cDNA | |
| | (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4: | |
| CTTC | ATGTTT CCAAAGTGCA TGAT | 24 |
| (2) | INFORMATION FOR SEQ ID NO:5: | |

Serial No.: 07/989,16

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 25 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single.
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: CDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

CTGGGCTTCA CTTCAGAGGA GAAAA

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- (2) INFORMATION FOR SEO ID NO:6:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 30 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

GCGGTACCCC AGCAGCCCGG CCTTGAAGAA

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- (2) INFORMATION FOR SEQ ID NO:7:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 30 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

GGGAATTCGC GGAGCCAGAC GGCACTGAAG

30

- (2) INFORMATION FOR SEQ ID NO:8:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: CDNA

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

- (2) INFORMATION FOR SEQ ID NO:9:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 25 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9: CAACTCATCA CCACTCTCTT CCATC
- (2) INFORMATION FOR SEQ ID NO:10:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 25 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE -TYPE: cDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:
 GCTGAGCCTA GCAGATTCAT GGCAC

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